ECE521:

Week 11, Lecture 20

27 March 2017: HMM learning/inference

With thanks to Russ Salakhutdinov

Examples of other perspectives

- Murphy 17.4
- End of Russell & Norvig 15.2

(Artificial Intelligence: A Modern Approach)

• Bishop 13.2.5, 8.4.4

Outline

- HMM learning and inference:
 - 1) Probability of an observed data sequence
 - 2) Learning the model parameters
 - 3) Inferring the most likely state sequence
- Thursday: Message passing
 - Introduction
 - Sum-product algorithm

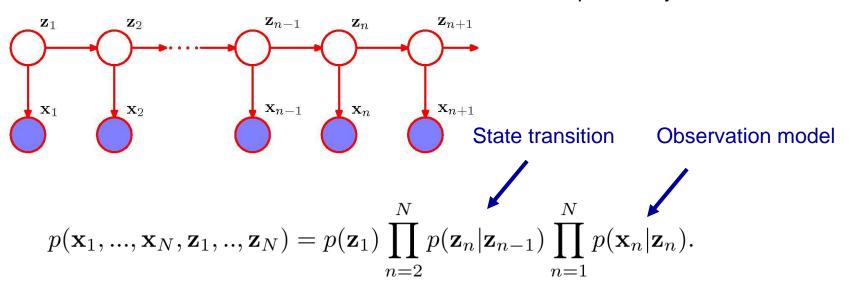
Recap of Hidden Markov Models

 First-order Markov chain generates hidden state sequence (known as transition probabilities):

$$p(\mathbf{z}_n = k | \mathbf{z}_{n-1} = j) = A_{jk}, \quad p(\mathbf{z}_1 = k) = \pi_k.$$

A set of output probability distributions (one per state) converts state path into sequence of observable symbols/vectors (known as emission probabilities): p(x_n|z_n, φ).

Can be e.g. Gaussian if **x** is continuous. Conditional probability table if **x** is discrete.



Recap of Hidden Markov Models

In the last lecture you looked at six examples:

- Sampling (generating states & observations)
- Prediction of the next state(s) given the current state (Example 2)
- Inferring the latent states behind an observed sequence (Example 6)

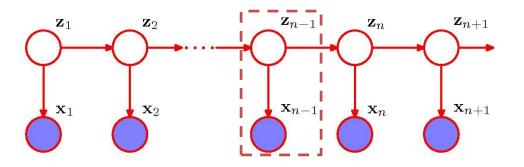
We build on Example 6 today

Three problems

• The joint distribution over the observed- and latent variables is given by:

$$p(\mathbf{X}, \mathbf{Z}|\theta) = p(\mathbf{z}_1|\boldsymbol{\pi}) \prod_{n=2}^{N} p(\mathbf{z}_n | \mathbf{z}_{n-1}, A) \prod_{n=1}^{N} p(\mathbf{x}_n | \mathbf{z}_n, \boldsymbol{\phi}),$$

where $\theta = \{\pi, A, \phi\}$ are the model parameters.



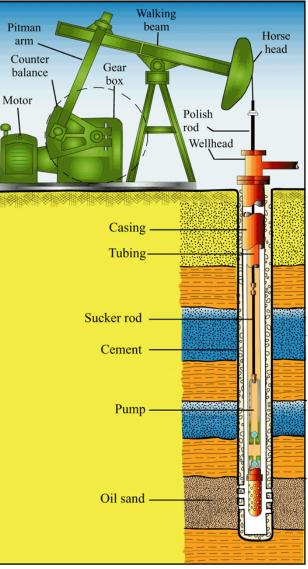
Three problems and three solutions:

- 1. Computing probabilities of observed sequences: Forward-backward algorithm
- 2. Learning of parameters: Baum-Welch algorithm
- 3. Inference of hidden state sequences: Viterbi algorithm +

Viterbi algorithm application: Casing Running

www.youtube.com/watch?v=F-HrLO5m_-s





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Maximum Likelihood for the HMM

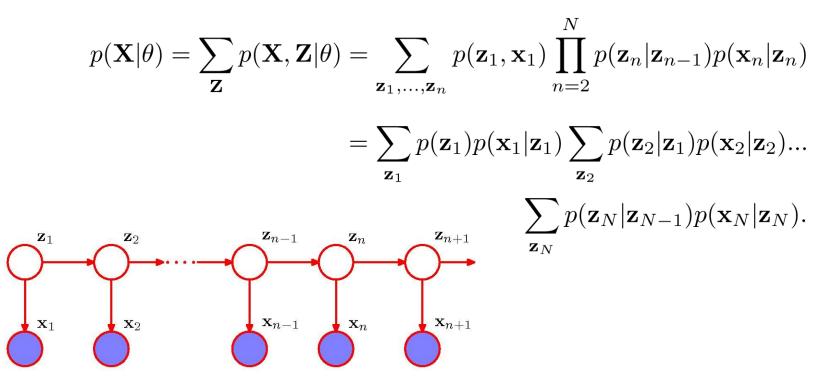
- We observe a dataset $\mathbf{X} = {\mathbf{x}_1, \dots, \mathbf{x}_N}$.
- The goal is to determine model parameters $\theta = \{\pi, A, \phi\}.$
- The probability of an observed sequence takes the form:

$$p(\mathbf{X}|\theta) = \sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\theta).$$
$$p(\text{observed sequence}) = \sum_{\text{all paths}} p(\text{observed outputs, state paths}).$$

- Recall that, in contrast to mixture models, the joint distribution p(X,Z | μ) does not factorize over n.
- It looks hard: N variables, each of which has K states. Hence K^N total paths.

Probability of an Observed Sequence

• Recalling slides 22-23 from Thursday (lecture 19), probabilities factorize:



• Dynamic Programming: By moving the summations inside, we can save a lot of work.

EM algorithm for HMMs: overview

• We cannot perform direct maximization (no closed-form solution):

$$p(\mathbf{X}|\theta) = \sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\theta).$$

- EM algorithm: we will derive an efficient algorithm for maximizing the likelihood function in HMMs (and, later, in linear state-space models).
- E-step: Compute the posterior distribution over latent variables:

$$p(\mathbf{Z}|\mathbf{X}, \theta^{old}).$$

• M-step: Maximize the expected complete data log-likelihood:

$$Q(\theta, \theta^{old}) = \sum_{\mathbf{Z}} p(\mathbf{Z} | \mathbf{X}, \theta^{old}) \log p(\mathbf{X}, \mathbf{Z} | \theta).$$

- If we knew the true state path, then ML parameter estimation would be trivial.
- We will first look at the E-step: Computing the true posterior distribution over the state paths.

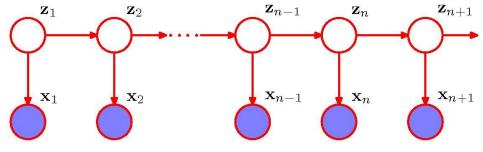
Inference of Hidden States

• We want to estimate the hidden states given observations. To start with, let us estimate a single hidden state:

$$\gamma(\mathbf{z}_n) = p(\mathbf{z}_n | \mathbf{X}) = \frac{p(\mathbf{X} | \mathbf{z}_n) p(\mathbf{z}_n)}{p(\mathbf{X})}$$

• Using the conditional-independence property, we obtain:

$$p(\mathbf{z}_n | \mathbf{X}) = \frac{p(\mathbf{x}_1, ..., \mathbf{x}_n | \mathbf{z}_n) p(\mathbf{x}_{n+1}, ..., \mathbf{x}_N | \mathbf{z}_n) p(\mathbf{z}_n)}{p(\mathbf{X})}$$
$$= \frac{p(\mathbf{x}_1, ..., \mathbf{x}_n, \mathbf{z}_n) p(\mathbf{x}_{n+1}, ..., \mathbf{x}_N | \mathbf{z}_n)}{p(\mathbf{X})} = \frac{\alpha(\mathbf{z}_n) \beta(\mathbf{z}_n)}{p(\mathbf{X})}$$



Inference of Hidden States

• Hence:

$$\gamma(\mathbf{z}_n) = \frac{p(\mathbf{x}_1, ..., \mathbf{x}_n, \mathbf{z}_n) p(\mathbf{x}_{n+1}, ..., \mathbf{x}_N | \mathbf{z}_n)}{p(\mathbf{X})} = \frac{\alpha(\mathbf{z}_n) \beta(\mathbf{z}_n)}{p(\mathbf{X})}.$$

 $\alpha(\mathbf{z}_n) \equiv p(\mathbf{x}_1, .., \mathbf{x}_n, \mathbf{z}_n)$

$$\beta(\mathbf{z}_n) \equiv p(\mathbf{x}_{n+1}, .., \mathbf{x}_N | \mathbf{z}_n). \blacktriangleleft$$

The joint probability of observing all of the data up to time n, and \mathbf{z}_n .

The conditional probability of all future data from time n+1 to N.

- Each α(z_n) and β(z_n) represent a set of K numbers, one for each of the possible settings of the 1-of-K binary vector z_n.
- We will derive an efficient recursive algorithm, known as the alpha-beta recursion, or forward-backward algorithm.

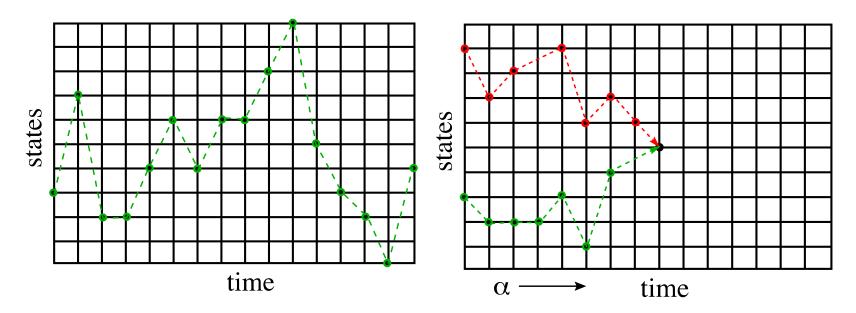
The Forward (a) Recursion

• The forward recursion:

• This enables us to easily (cheaply) compute the desired likelihood.

The Forward (α) Recursion

• The forward recursion:



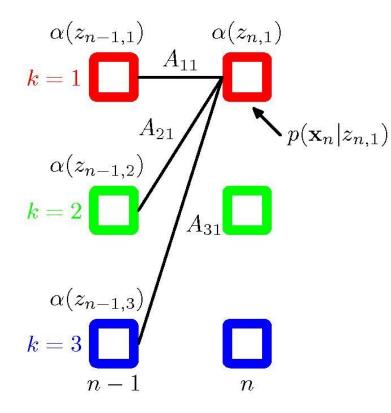
Exponentially many paths.

At each node, sum up the values of all incoming paths.

• This is exactly dynamic programming.

The Forward (α) Recursion

• Illustration of the forward recursion



Here $\alpha(z_{n,1})$ is obtained by:

- Taking the elements $\alpha(z_{n-1}, j)$
- Summing them up with weights A_{j1} , corresponding to $p(\mathbf{z}_n | \mathbf{z}_{n-1})$
- Multiplying by the data contribution $p(\mathbf{x}_n | \mathbf{z}_{n,1})$.

$$\alpha(\mathbf{z}_n) = p(\mathbf{x}_n | \mathbf{z}_n) \sum_{\mathbf{z}_{n-1}} \alpha(\mathbf{z}_{n-1}) p(\mathbf{z}_n | \mathbf{z}_{n-1})$$

• The initial condition is given by: $\alpha(\mathbf{z}_1) = p(\mathbf{x}_1 | \mathbf{z}_1) p(\mathbf{z}_1) = \prod_{k=1}^K \left[\pi_k p(\mathbf{x}_1 | \boldsymbol{\phi}_k) \right]^{z_{1k}}.$

The Backward (β) Recursion

• There is also a simple recursion for $\beta(\mathbf{z}_n)$:

$$\beta(\mathbf{z}_{n}) = p(\mathbf{x}_{n+1}, ..., \mathbf{x}_{N} | \mathbf{z}_{n})$$

$$= \sum_{\mathbf{z}_{n+1}} p(\mathbf{x}_{n+1}, ..., \mathbf{x}_{N}, \mathbf{z}_{n+1} | \mathbf{z}_{n})$$

$$= \sum_{\mathbf{z}_{n+1}} p(\mathbf{x}_{n+1}, ..., \mathbf{x}_{N} | \mathbf{z}_{n+1}, \mathbf{z}_{n}) p(\mathbf{z}_{n+1} | \mathbf{z}_{n})$$

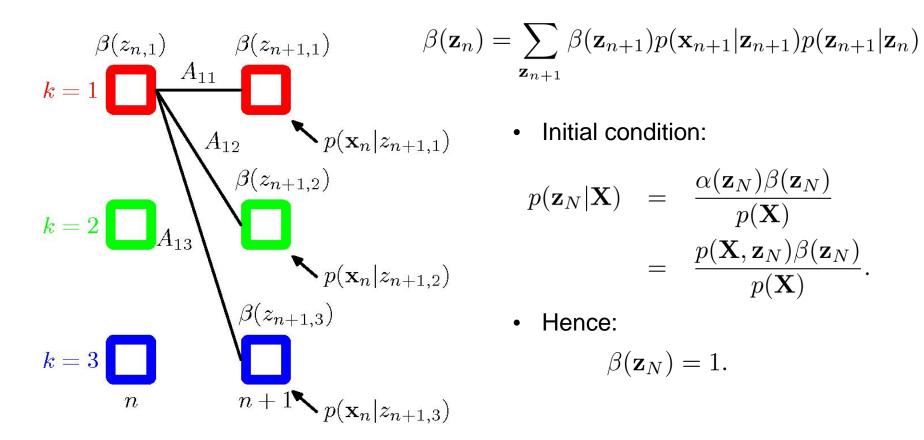
$$= \sum_{\mathbf{z}_{n+1}} p(\mathbf{x}_{n+1}, ..., \mathbf{x}_{N} | \mathbf{z}_{n+1}) p(\mathbf{z}_{n+1} | \mathbf{z}_{n})$$

$$= \sum_{\mathbf{z}_{n+1}} p(\mathbf{x}_{n+2}, ..., \mathbf{x}_{N} | \mathbf{z}_{n+1}) p(\mathbf{x}_{n+1} | \mathbf{z}_{n+1}) p(\mathbf{z}_{n+1} | \mathbf{z}_{n})$$

$$= \sum_{\mathbf{z}_{n+1}} \beta(\mathbf{z}_{n+1}) p(\mathbf{x}_{n+1} | \mathbf{z}_{n+1}) p(\mathbf{z}_{n+1} | \mathbf{z}_{n})$$

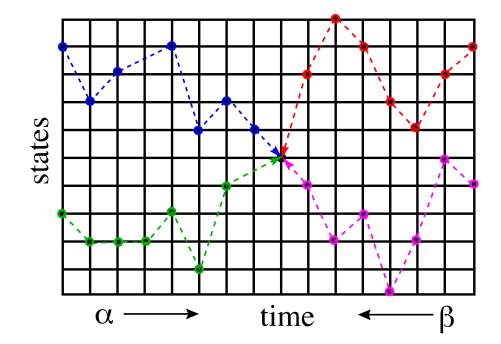
The Backward (β) Recursion

Illustration of the backward recursion



The Backward (β) Recursion

- $\alpha(z_{nk})$ gives total inflow of probability to node (n,k).
- $\beta(z_{nk})$ gives total outflow of probability.



 In fact, we can do one forward pass to compute all the *α*(**z**_n) and one backward pass to compute all the β(**z**_n) and then compute any γ(**z**_n) we want. Total cost is **O**(K²N).

Computing Likelihood

Note that

$$\sum_{\mathbf{z}_n} \gamma(\mathbf{z}_n) = \sum_{\mathbf{z}_n} p(\mathbf{z}_n | \mathbf{X}) = 1.$$

• We can compute the likelihood at any time using α - β recursion:

$$p(\mathbf{X}|\theta) = \sum_{\mathbf{z}_n} \alpha(\mathbf{z}_n) \beta(\mathbf{z}_n).$$

 In the forward calculation we proposed originally, we did this at the final time step n = N.

$$p(\mathbf{X}|\theta) = \sum_{\mathbf{z}_N} \alpha(\mathbf{z}_N).$$

because $\beta(\mathbf{z}_n)=1$.

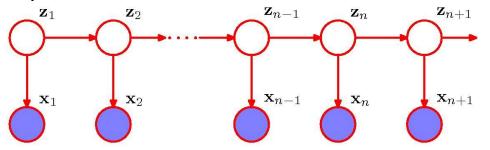
Monitoring this quantity is a good way to check for convergence during EM

Two-Frame Inference

• We will also need the cross-time statistics for adjacent time steps:

$$\begin{aligned} \xi(\mathbf{z}_{n-1}, \mathbf{z}_n) &= p(\mathbf{z}_{n-1}, \mathbf{z}_n | \mathbf{X}) \\ &= \frac{p(\mathbf{X} | \mathbf{z}_{n-1}, \mathbf{z}_n) p(\mathbf{z}_{n-1}, \mathbf{z}_n)}{p(\mathbf{X})} \\ &= \frac{p(\mathbf{x}_1, \dots, \mathbf{x}_{n-1} | \mathbf{z}_{n-1}) p(\mathbf{x}_n | \mathbf{z}_n) p(\mathbf{x}_{n+1}, \dots, \mathbf{x}_N | \mathbf{z}_n) p(\mathbf{z}_n | \mathbf{z}_{n-1}) p(\mathbf{z}_{n-1})}{p(\mathbf{X})} \\ &= \frac{\alpha(\mathbf{z}_{n-1}) p(\mathbf{x}_n | \mathbf{z}_n) p(\mathbf{z}_n | \mathbf{z}_{n-1}) \beta(\mathbf{z}_n)}{p(\mathbf{X})}. \end{aligned}$$

- This is a K x K matrix with elements (*i*,*j*) representing the expected number of transitions from state *i* to state *j* that begin at time *n*-1, given all the observations.
- Whereas γ is the marginal posterior distribution of a latent variable, ξ is the joint posterior distribution of two successive latent variables



It can be computed with the same α- and β recursions.

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The second problem: EM algorithm

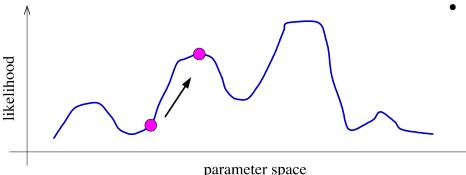
- Intuition: if only we knew the true state path then ML parameter estimation would be trivial.
- E-step: Compute the posterior distribution over the state path using α-β recursion (dynamic programming):

$$p(\mathbf{Z}|\mathbf{X}, \theta^{old}).$$

 M-step: Maximize the expected complete data log-likelihood (parameter reestimation):

$$Q(\theta, \theta^{old}) = \sum_{\mathbf{Z}} p(\mathbf{Z} | \mathbf{X}, \theta^{old}) \log p(\mathbf{X}, \mathbf{Z} | \theta).$$

• We then iterate. This is also known as a **Baum-Welch algorithm** (that is, EM applied to HMMs).



• In general, finding the ML parameters is NP hard, so initial conditions matter a lot

Complete Data Log-likelihood

• The complete data log-likelihood takes the form:

$$\log p(\mathbf{X}, \mathbf{Z} | \theta) = \log \left[p(\mathbf{z}_1 | \boldsymbol{\pi}) \prod_{n=2}^{N} p(\mathbf{z}_n | \mathbf{z}_{n-1}, A) \prod_{n=1}^{N} p(\mathbf{x}_n | \mathbf{z}_n, \phi) \right]$$

$$= \log \left[\prod_{k=1}^{K} \pi_k^{z_{1k}} \prod_{n=2}^{N} \prod_{k=1}^{K} \prod_{j=1}^{K} A_{jk}^{z_{n-1,j}, z_{nk}} \prod_{n=1}^{N} \prod_{k=1}^{K} p(\mathbf{x}_n | \mathbf{z}_n)^{z_{nk}} \right]$$

$$= \sum_{k=1}^{K} z_{1k} \log \pi_k + \sum_{n=2}^{N} \sum_{k=1}^{K} \sum_{j=1}^{K} [z_{n-1,j} z_{nk}] \log A_{jk} + \sum_{n=1}^{N} \sum_{k=1}^{K} z_{nk} \log p(\mathbf{x}_n | \mathbf{z}_n).$$
transition model observation model

Expected Complete Data Log-likelihood

• The complete data log-likelihood takes the form:

$$Q(\theta, \theta^{old}) = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \theta^{old}) \log p(\mathbf{X}, \mathbf{Z}|\theta).$$
$$= \sum_{k=1}^{K} \gamma(z_{1k}) \log \pi_k + \sum_{n=2}^{N} \sum_{k=1}^{K} \sum_{j=1}^{K} \xi(z_{n-1,j}z_{nk}) \log A_{jk} + \sum_{n=1}^{N} \sum_{k=1}^{K} \gamma_{nk} \log p(\mathbf{x}_n|\mathbf{z}_n).$$

• Recall that in the E-step, we evaluate:

$$\gamma(\mathbf{z}_n) = p(\mathbf{z}_n | \mathbf{X}).$$

$$\xi(\mathbf{z}_{n-1}, \mathbf{z}_n) = p(\mathbf{z}_{n-1}, \mathbf{z}_n | \mathbf{X}).$$

• In the M-step, we optimize Q with respect to parameters $\theta = \{\pi, A, \phi\}$.

Parameter Estimation: π and A

• Initial state distribution: Using Lagrange multipliers, the expected number of times in state k at time 1 is: $\gamma(z_{1k})$

$$\pi_k^{new} = \frac{\gamma(z_{1k})}{\sum_{j=1}^K \gamma(z_{1j})}.$$

• Expected number of transitions from state *j* to *k* which begin at time *n*-1:

$$\xi(\mathbf{z}_{n-1,j},\mathbf{z}_{n,k}) = p(\mathbf{z}_{n-1,j},\mathbf{z}_{n,k}|\mathbf{X}),$$

and the estimated transition probabilities work out to be:

$$A_{jk}^{new} = \frac{\sum_{n=2}^{N} \xi(z_{n-1,j}, z_{nk})}{\sum_{l=1}^{K} \sum_{n=2}^{N} \xi(z_{n-1,j}, z_{nl})}.$$

- The EM algorithm must be initialized by choosing starting values for π and A.
- Note that any elements of *π* or A that initially are set to zero will remain zero in subsequent EM updates.

Parameter Estimation: Emission Model

For the case of discrete multinomial observed variables, the observation model takes the form: DKSame as fitting Bernoulli

$$p(\mathbf{x}_n | \mathbf{z}_n, \boldsymbol{\phi}) = \prod_{i=1} \prod_{k=1} \mu_{ik}^{x_{ni} z_{nk}}.$$

- And the corresponding M-step update: μ_{ik}^{new}
- For the case of the Gaussian emission model: K $p(\mathbf{x}_n | \mathbf{z}_n, \boldsymbol{\phi}) = \prod \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)^{z_{nk}}.$ k=1
- And the corresponding M-step updates:

$$\boldsymbol{\mu}_{k}^{new} = \frac{1}{N_{k}} \sum_{n}^{n} \gamma(z_{nk}) \mathbf{x}_{n}, \quad N_{k} = \sum_{n}^{n} \gamma(z_{nk}), \quad \begin{array}{c} \text{barrie as futting a Gamma Matrix}\\ \text{mixture model.} \end{array}$$
$$\boldsymbol{\Sigma}_{k}^{new} = \frac{1}{N_{k}} \sum_{n=1}^{N} \gamma(z_{nk}) (\mathbf{x}_{n} - \boldsymbol{\mu}_{k}) (\mathbf{x}_{n} - \boldsymbol{\mu}_{k})^{T},$$

Remember:

$$\gamma(\mathbf{z}_n) = p(\mathbf{z}_n | \mathbf{X}).$$

Same as fitting a Gaussian

$$=\frac{\sum_{n=1}^{N}\gamma(z_{nk})x_{ni}}{\sum_{n=1}^{N}\gamma(z_{nk})}.$$

mixture model.

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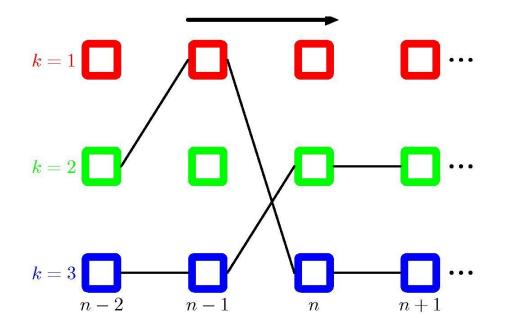
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The third problem: Viterbi Decoding

- The numbers γ(z_n) above gave the probability distribution over all states at any time.
- By choosing the state γ*(z_n) with the largest probability at each time, we can make an "average" state path. This is the path with the maximum expected number of correct states.
- To find the single best path, we do Viterbi decoding which is a dynamic programming algorithm applied to this problem.
- The recursions look the same, except with 'max' instead of Σ .
- Same dynamic programming trick: instead of summing, we keep the term with the highest value at each node.
- There is also a modified EM (Baum-Welch) training based on the Viterbi decoding. Like K-means instead of mixtures of Gaussians.

Viterbi Decoding

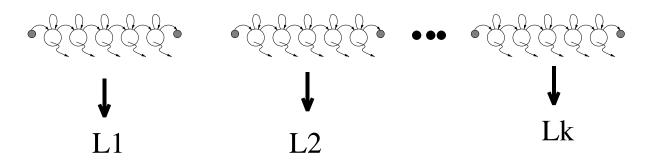
• A fragment of the HMM lattice showing two possible paths:



- Viterbi decoding efficiently determines the most probable path from the exponentially many possibilities.
- The probability of each path is given by the product of the elements of the transition matrix A_{jk} , along with the emission probabilities associated with each node in the path.

Using HMMs for Recognition

- We can use HMMs for recognition by:
 - Training one HMM for each class (requires labelled training data)
 - Evaluating the probability of an unknown sequence under each HMM
 - Classifying the unknown sequence by choosing an HMM with highest likelihood

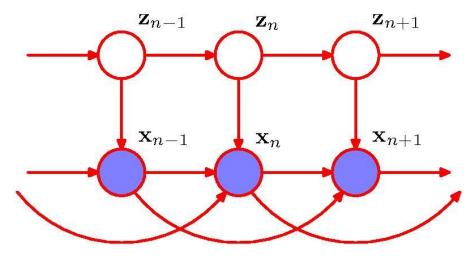


- This requires the solution of two problems:
 - Given a model, evaluate the probability of a sequence. (We can do this exactly and efficiently.)
 - Given some training sequences, estimate the model parameters. (We can find the local maximum using EM.)

Bonus

Autoregressive HMMs

• One limitation of the standard HMM is that it is poor at capturing longrange correlations between observations, as these have to be mediated via the first order Markov chain of hidden states.



- Autoregressive HMM: The distribution over x_n depends also on a subset of previous observations.
- The number of additional links must be limited to avoid an excessive number of free parameters.
- The graphical model framework motivates a number of different models based on HMMs.

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